

Amendments to the Claims:

This listing of claims replaces all prior versions and listings of claims in the application:

Listing of Claims:

1-14. (Withdrawn)

15. (Original) A kit for detecting dengue virus comprising:

A first dengue virus-specific primer, which is 18 to 28 nucleotides in length and includes at least 18 nucleotides of SEQ ID NO:1; and

A second dengue virus-specific primer, which is 18 to 28 nucleotides in length and includes at least 18 nucleotides of SEQ ID NO:2.

16. (Original) The kit of claim 15, further comprising a known amount of a competitor nucleic acid with length detectably different from the dengue virus RNA.

17. (Original) The kit of claim 15, wherein the first dengue virus-specific primer is 18 to 23 nucleotides in length.

18. (Original) The kit of claim 15, wherein the first dengue virus-specific primer is the nucleotide sequence of SEQ ID NO:1.

19. (Original) The kit of claim 15, wherein the second dengue virus-specific primer is 18 to 23 nucleotides in length.

20. (Original) The kit of claim 15, wherein the second dengue virus-specific primer is the nucleotide sequence of SEQ ID NO:2.

21. (Original) The kit of claim 17, wherein the second dengue virus-specific primer is 18 to 23 nucleotides in length.

22. (Original) The kit of claim 18, wherein the second dengue virus-specific primer is the nucleotide sequence of SEQ ID NO:2.

23. (Original) A nucleic acid, which is 18 to 28 nucleotides in length and includes at least 18 consecutive nucleotides of SEQ ID NO:1.

24. (Original) The nucleic acid of claim 23, wherein the nucleic acid is 18 to 23 nucleotides in length and includes at least 18 consecutive nucleotides of SEQ ID NO:1.

25. (Original) The nucleic acid of claim 23, wherein the nucleic acid is the nucleotide sequence of SEQ ID NO:1.

26. (Currently Amended) A nucleic acid, which is 18 to ~~28~~ 25 nucleotides in length and includes at least 18 consecutive nucleotides of SEQ ID NO:2.

27. (Original) The nucleic acid of claim 26, wherein the nucleic acid is 18 to 23 nucleotides in length and includes at least 18 consecutive nucleotides of SEQ ID NO:2.

28. (Currently Amended) A ~~The nucleic acid of claim 26~~ nucleic acid, wherein the nucleic acid is the nucleotide sequence of SEQ ID NO:2.

29. (Original) An isolated nucleic acid comprising a fragment of a dengue viral genome or a DNA copy thereof, wherein the fragment includes:

a first sequence that is complementary or identical to at least 18 consecutive nucleotides of SEQ ID NO:1;

a second sequence that is complementary or identical to at least 18 consecutive nucleotides of SEQ ID NO:2; and

a non-naturally occurring deletion or insertion, the deletion or insertion occurring in a region of the fragment flanked by the first and the second sequence.

30. (Original) The nucleic acid of claim 29, wherein the first sequence is complementary or identical to SEQ ID NO:1 and the second sequence that is complementary or identical to SEQ ID NO:2.

31. (New) The nucleic acid of claim 29, wherein the nucleic acid consists of:
a first sequence that is complementary or identical to at least 18 consecutive nucleotides of SEQ ID NO:1;

a second sequence that is complementary or identical to at least 18 consecutive nucleotides of SEQ ID NO:2; and

a third sequence that is flanked by the first and second sequence, wherein the third sequence is complementary or identical to a genomic Dengue virus region that is (a) naturally flanked by the first and the second sequence and (b) includes a non-naturally occurring deletion or insertion.

32. (New) The nucleic acid of claim 31, wherein the first sequence is complementary or identical to SEQ ID NO:1 and the second sequence is complementary or identical to SEQ ID NO:2.